GenCore version 5.1.3

OM nucleic - nucleic search, using sw model

Run on: February 16, 2003, 22:05:25 ; Search time 51.0802 Seconds
(without alignments)
13999.354 Million cell updates/sec

Title: US-09-497-967-3
Perfect score: 1404
Sequence: 1 atgaaaaataatattttagt.....tgatttcttattatttatta 1404

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 424239 seqs, 254661826 residues

Total number of hits satisfying chosen parameters: 848478

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Published_Applications_NA:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	Result No.	Score	% Query Match	Length	DB	ID	Description
c	1	60	4.3	1635	10	US-09-864-761-20241	Sequence 20241, A
c	2	60	4.3	1973	10	US-09-864-761-3471	Sequence 3471, Ap
c	3	52.2	3.7	510	10	US-09-864-761-18737	Sequence 18737, A
	4	46.6	3.3	684973	10	US-09-263-959-1	Sequence 1, Appli
	5	45.8	3.3	1075	10	US-09-864-761-19241	Sequence 19241, A
	6	45.8	3.3	1403	10	US-09-864-761-2513	Sequence 2513, Ap
c	7	43.4	3.1	2120	10	US-09-798-042-95	Sequence 95, Appl
	8	43.4	3.1	2129	10	US-09-159-469-39	Sequence 39, Appl
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	18	40	2.8	250	10	US-09-815-242-3253	Sequence 3253, Ap
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c	22	40	2.8	1476	10	US-09-815-242-8394	Sequence 8394, Ap
	23	40	2.8	1908	9	US-09-286-488-50	Sequence 50, Appl
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	27	40	2.8	3402	10	US-09-737-178-86	Sequence 86, Appl
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	38	39.2	2.8	155074	9	US-10-026-188-6	Sequence 6, Appli
	39	39	2.8	4634	10	US-09-995-587A-10	Sequence 10, Appl
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	45	38.6	2.7	513509	9	US-09-754-853A-4	Sequence 4, Appli

ALIGNMENTS

RESULT 1
US-09-864-761-20241/c
; Sequence 20241, Application US/09864761
; Patent No. US20020048763A1
; GENERAL INFORMATION:
; APPLICANT: Penn, Sharron G.
; APPLICANT: Rank, David R.
; APPLICANT: Hanzel, David K.
; APPLICANT: Chen, Wensheng
; TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FO
; TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
; FILE REFERENCE: Aeomica-X-1
; CURRENT APPLICATION NUMBER: US/09/864,761
; CURRENT FILING DATE: 2001-05-23
; PRIOR APPLICATION NUMBER: US 60/180,312
; PRIOR FILING DATE: 2000-02-04
; PRIOR APPLICATION NUMBER: US 60/207,456
; PRIOR FILING DATE: 2000-05-26
; PRIOR APPLICATION NUMBER: US 09/632,366
; PRIOR FILING DATE: 2000-08-03
; PRIOR APPLICATION NUMBER: GB 24263.6
; PRIOR FILING DATE: 2000-10-04
; PRIOR APPLICATION NUMBER: US 60/236,359
; PRIOR FILING DATE: 2000-09-27
; PRIOR APPLICATION NUMBER: PCT/US01/00666
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00667
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00664
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00669
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00665
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00668
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00663
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00662
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00661
; PRIOR FILING DATE: 2001-01-30

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;  PRIOR APPLICATION NUMBER: PCT/US01/00670
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: US 60/234,687
;  PRIOR FILING DATE: 2000-09-21
;  PRIOR APPLICATION NUMBER: US 09/608,408
;  PRIOR FILING DATE: 2000-06-30
;  PRIOR APPLICATION NUMBER: US 09/774,203
;  PRIOR FILING DATE: 2001-01-29
;  NUMBER OF SEQ ID NOS: 49117
;  SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
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;   LENGTH: 1635
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: MAP TO AC006547.9
;   OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL =  17
;   OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL =  11
;   OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 8.6
;   OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL =  9
;   OTHER INFORMATION: EXPRESSED IN HBL100, SIGNAL =  12
;   OTHER INFORMATION: EXPRESSED IN BT474, SIGNAL =  11
;   OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL =  10
;   OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 8.4
;   OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL =  10
;   OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL =  12
US-09-864-761-20241

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  Best Local Similarity   41.5%;  Pred. No. 4.5e-05;
  Matches  463;  Conservative    0;  Mismatches  650;  Indels    3;  Gaps    1;

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Qy     120 TCCTGCAAATTGTGTTAATTGTTAGAAAAACTTTTATTATAATAATGCTGCTGCTTTCGT 179
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Db    1310 TGGTGGTAGTGGTGATGGTGGTAGTGATGGTGGTTGTAGTGATGGTGGTGGTGATGGTGG 1251

Qy     180 TCCTGGTGCTAGTACGTGTACACCTTGTCCATAAAAAAAAGATGCTGGTGCTTAACCAAA 239
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Db    1250 TGATGGTGGTAGTGATGGTGGTGGTGGTGGTGGTAGTGATGGTGATGGTGGTGGTGGTGA 1191

Qy     240 TCCACCTGCTACTGCTAATTTAGTCACATAATGTAACGTTAAATGCCCTGCTGGTACCGC 299
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Qy     477 ATAATGTAACGTCGCATGTCCTACTGGTACTGCACTTGATGATGGAGTAACTACTGATTA 536
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Db     950 TGATGGTGATGGTGGTGGTGGTGGTAGTGATGGTGGTGGTGGTGGTGGTGGTAGTGATGG 891

Qy     537 TGTTAGATCATTCACAGAATGTGTTAAATGTAGACTTAACTTTTACTATAATGGTAATAA 596
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Db     890 TGGTGGTGGTGGTAGTGATGGTGGTGGTGGTGGTGGTAGTGATGGTGGTGGTGGTGGTGG 831

Qy     597 TGGTAATACTCCTTTCAATCCAGGTAAAAGTTAATGCACACCTTGTCCGGCAATTAAACC 656
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Db     830 TGGTAGTGATGGTGGTGGTGGTGGTGATGGTGGTGGTGGTAGTGATGGTGGTGGTGGTGG 771

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Qy     777 TGAATGTACTAATTGTGCTCCTAACTTTTACAATAATAATGCTCCTAATTTCAATCCAGG 836
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Qy     897 TGGTGCCGCTACTTTAGCCAAATAATGTAATATTGCATGCCCTGATGGTACTGCAATTGC 956
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RESULT 2
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; Sequence 3471, Application US/09864761
; Patent No. US20020048763A1
; GENERAL INFORMATION:
;  APPLICANT: Penn, Sharron G.
;  APPLICANT:  Rank, David R.
;  APPLICANT:  Hanzel, David K.
;  APPLICANT:  Chen, Wensheng
;  TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FO
;  TITLE OF INVENTION:  GENE EXPRESSION ANALYSIS BY MICROARRAY
;  FILE REFERENCE: Aeomica-X-1
;  CURRENT APPLICATION NUMBER: US/09/864,761
;  CURRENT FILING DATE:  2001-05-23
;  PRIOR APPLICATION NUMBER: US 60/180,312
;  PRIOR FILING DATE: 2000-02-04
;  PRIOR APPLICATION NUMBER: US 60/207,456
;  PRIOR FILING DATE: 2000-05-26
;  PRIOR APPLICATION NUMBER: US 09/632,366
;  PRIOR FILING DATE: 2000-08-03
;  PRIOR APPLICATION NUMBER: GB 24263.6
;  PRIOR FILING DATE: 2000-10-04
;  PRIOR APPLICATION NUMBER: US 60/236,359
;  PRIOR FILING DATE: 2000-09-27
;  PRIOR APPLICATION NUMBER: PCT/US01/00666
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00667
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00664
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00669
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00665
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00668
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00663
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;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00662
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00661
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00670
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: US 60/234,687
;  PRIOR FILING DATE: 2000-09-21
;  PRIOR APPLICATION NUMBER: US 09/608,408
;  PRIOR FILING DATE: 2000-06-30
;  PRIOR APPLICATION NUMBER: US 09/774,203
;  PRIOR FILING DATE: 2001-01-29
;  NUMBER OF SEQ ID NOS: 49117
;  SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
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;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: MAP TO AC006547.9
;   OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL =  17
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;   OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL =  10
;   OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 8.4
;   OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL =  10
;   OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL =  12
US-09-864-761-3471

  Query Match             4.3%;  Score 60;  DB 10;  Length 1973;
  Best Local Similarity   41.5%;  Pred. No. 4.9e-05;
  Matches  463;  Conservative    0;  Mismatches  650;  Indels    3;  Gaps    1;

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Qy     180 TCCTGGTGCTAGTACGTGTACACCTTGTCCATAAAAAAAAGATGCTGGTGCTTAACCAAA 239
           |  ||||| ||||    ||     | ||       |   | | || ||||| |       |
Db    1404 TGATGGTGGTAGTGATGGTGGTGGTGGTGGTGGTAGTGATGGTGATGGTGGTGGTGGTGA 1345

Qy     240 TCCACCTGCTACTGCTAATTTAGTCACATAATGTAACGTTAAATGCCCTGCTGGTACCGC 299
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Db    1344 TGGTGGTGGTGGTGGTGATGGTGGTAGTGATGGTGATGGTGGTGGTGGTAGTGATGG 1285

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Qy     360 TAATTTTTATAATGAAAATGCTCCAAATTTTAATGCAGGTGCTAGTACATGCACAGCTTG 419
           || |  |  |  ||    || |     |  || ||  |||| | ||    |    | | |
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Db     444 TGGTGATGGTGGTGGTGGTGGTGGTGGTAGTGATGG 409


RESULT 3
US-09-864-761-18737/c
; Sequence 18737, Application US/09864761
; Patent No. US20020048763A1
; GENERAL INFORMATION:
;  APPLICANT: Penn, Sharron G.
;  APPLICANT:  Rank, David R.
;  APPLICANT:  Hanzel, David K.
;  APPLICANT:  Chen, Wensheng
;  TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FO
;  TITLE OF INVENTION:  GENE EXPRESSION ANALYSIS BY MICROARRAY
;  FILE REFERENCE: Aeomica-X-1
;  CURRENT APPLICATION NUMBER: US/09/864,761
;  CURRENT FILING DATE:  2001-05-23
;  PRIOR APPLICATION NUMBER: US 60/180,312
;  PRIOR FILING DATE: 2000-02-04
;  PRIOR APPLICATION NUMBER: US 60/207,456
;  PRIOR FILING DATE: 2000-05-26
;  PRIOR APPLICATION NUMBER: US 09/632,366
;  PRIOR FILING DATE: 2000-08-03
;  PRIOR APPLICATION NUMBER: GB 24263.6
;  PRIOR FILING DATE: 2000-10-04
;  PRIOR APPLICATION NUMBER: US 60/236,359
;  PRIOR FILING DATE: 2000-09-27
;  PRIOR APPLICATION NUMBER: PCT/US01/00666
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00667
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00664
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00669
;  PRIOR FILING DATE: 2001-01-30
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;  PRIOR APPLICATION NUMBER: PCT/US01/00665
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00668
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00663
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00662
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00661
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00670
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: US 60/234,687
;  PRIOR FILING DATE: 2000-09-21
;  PRIOR APPLICATION NUMBER: US 09/608,408
;  PRIOR FILING DATE: 2000-06-30
;  PRIOR APPLICATION NUMBER: US 09/774,203
;  PRIOR FILING DATE: 2001-01-29
;  NUMBER OF SEQ ID NOS: 49117
;  SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
; SEQ ID NO 18737
;   LENGTH: 510
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: MAP TO AL035457.11
;   OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 1.2
;   OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 0.85
;   OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 1.2
;   OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 0.9
;   OTHER INFORMATION: EXPRESSED IN BT474, SIGNAL = 4.6
;   OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 0.93
;   OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 0.95
;   OTHER INFORMATION: NT HIT: U01287.1, EVALUE 1.30e+00
US-09-864-761-18737

  Query Match             3.7%;  Score 52.2;  DB 10;  Length 510;
  Best Local Similarity   46.2%;  Pred. No. 0.002;
  Matches  174;  Conservative    0;  Mismatches  203;  Indels    0;  Gaps    0;

Qy    162 TAATGCTGCTGCTTTCGTTCCTGGTGCTAGTACGTGTACACCTTGTCCATAAAAAAAAGA 221
          ||  | |||||     | |  || ||| || |    | |  |  |      |     |||
Db    491 TACCGATGCTGAAGCAGGTGTTGATGCAGGTGCTGAAAAAGGTGTTGGTGCAGGTGCAGA 432

Qy    222 TGCTGGTGCTTAACCAAATCCACCTGCTACTGCTAATTTAGTCACATAATGTAACGTTAA 281
          |||||  ||||       ||| |   |||    ||| ||   |   ||     |   |||
Db    431 TGCTGATGCTGGTAGTAATGCTGATGCAGGCGCTGATGCTGATGCAGGTATTGGTGTTGG 372

Qy    282 ATGCCCTGCTGGTACCGCAATTGCAGGTGGAGCAACAGATTATGCAGCAATAATCACAGA 341
              || || | | |   ||| ||  || ||   | |   ||  |    |    ||||
Db    371 TGCAGATGTTGATGCAGATGTTGAAGCAGGTGCTGAAAAAGGTGTTGATGCAGGTGCAGA 312

Qy    342 ATGTGTTAATTGTAGAATTAATTTTTATAATGAAAATGCTCCAAATTTTAATGCAGGTGC 401
            ||  |  ||||   |  |  |  || |||   |  | || || || |||||| ||
Db    311 TGCTGATGCTGGTAGTGATGCTGATGGTAGTGATGCTTGTGCAGATGTTGATGCAGATGT 252

Qy    402 TAGTACATGCACAGCTTGTCCGGTAAACAGAGTTGGTGGTGCATTGACTGCTGGTAATGC 461
          |    || |    | |       |  | | |  || || ||||    | | || | ||||
Db    251 TGAAGCAGGTGTTGATGCAGGTGCTGAAAAATGTGTTGATGCAAGTGCAGATGCTGATGC 192

Qy    462 CGCTACCATAGTCGCATAATGTAACGTCGCATGTCCTACTGGTACTGCACTTGATGATGG 521
            | ||     |  |       ||| |  ||  || |   || |  |||| || ||| |
Db    191 TGGTAGTGATGCTGATGCTGGTAGTGATGCTGGTGCAGATGTTGATGCAGATGTTGAAGC 132

Qy    522 AGTAACTACTGATTATG 538
          |    |  || | |||
Db    131 GGGTGTTGATGCTGATG 115


RESULT 4
US-09-263-959-1
; Sequence 1, Application US/09263959
; Patent No. US20020150891A1
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;  GENERAL INFORMATION:
;    APPLICANT:  Hood, Leroy E.
;    APPLICANT:  Rowen, Lee
;    APPLICANT:  Koop, Ben F.
;    TITLE OF INVENTION:  DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH U
;    NUMBER OF SEQUENCES:  1279
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Seed and Berry LLP
;      STREET:  6300 Columbia Center, 701 Fifth Avenue
;      CITY:  Seattle
;      STATE:  Washington
;      COUNTRY:  US
;      ZIP:  98104-7092
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/263,959
;      FILING DATE:  05-MAR-1999
;      CLASSIFICATION:
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  McMasters, David D.
;      REGISTRATION NUMBER:  33,963
;      REFERENCE/DOCKET NUMBER:  920010.426C2
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (206) 622-4900
;      TELEFAX:  (206) 682-6031
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  684973 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
US-09-263-959-1

  Query Match             3.3%;  Score 46.6;  DB 10;  Length 684973;
  Best Local Similarity   42.6%;  Pred. No. 1.1;
  Matches  363;  Conservative    0;  Mismatches  484;  Indels    6;  Gaps    2;

Qy    160 AATAATGCTGCTGCTTTCGTTCCTGGTGCTAGTACGTGTACACCTTGTCCATAAAAAAAA 219
          | || |  || || |     |  || | ||  | |   |||| ||   || |    || |
Db  12747 AGTACTATTGGTGTTACAACTAATGCTACTGTTCCCAATACAACTGCCCCTTTCCCAACA 12806

Qy    220 GATGCTGGTGCTTAACCAAATCCACCTGCTACTGCTAATTTAGTCACATAATGTAACGTT 279
           |||||  || ||       | |    || ||  | |||    |  |||||  | |
Db  12807 AATGCTAGTACTGCTAGCACTAATGCTACTGTTCCTATCACAACCACATGTTTTGCAACA 12866

Qy    280 AAATGCCCTGCTGGTACCGCAATTGCAGGTGGAGCAACAGATTATGCAGCAATAATCACA 339
          |       || || |||  | | |||   ||   |         |||  |  |    |||
Db  12867 AGTACTATTGGTGTTACAACTAATGCTACTGTTCCCGATACAACTGCCCCTTTCCCAACA 12926

Qy    340 GAATGTGTTAATTGTAGAATTAATTTTTATAATGAAAATGCTCCAAATTTTAATGCAGGT 399
          |   |  || |  ||| | ||||  |  || |   |   | | |    | ||||
Db  12927 AATACTACTACTGCTAGCACTAATGCTACTATTCCTATCACAACCACACCTTTTGCA--- 12983

Qy    400 GCTAGTACATGCACAGCTTGTCCGGTAAACAGAGTTGGTGGTGCATTGACTGCTGGTAAT 459
           | |||||  |  | |  |     |   |  |||  || | ||    || ||      |
Db  12984 ACAAGTACTATTAGTGTTACAACTAGTACTACTGTTCCTGATACAACTGCTCCTTTCCCT 13043

Qy    460 GCCGCTACCATAGTCGCATAATGTAACGTCGCATGTCCTACTGGTACTGCACTTGATGAT 519
           |  ||| |     ||     ||| |  |   |  | ||   ||  |       | |
Db  13044 ACAAGTACTACTAGTGCTAGCACTAATGCTACCCCTGTTCCTATCACAACCACACTTTTT 13103

Qy    520 GGAGTAACTACTGATTATGTTAGATCATTCACAGAATGTGTTAAATGTAGACTTAACTTT 579
          | |  || |||| |  ||||| | |    ||      ||||     ||| |  |   |
Db  13104 GCAACAAGTACTATTGGTGTTACAACTGGTAC---TACTGTTCCTGATACAACTGCTCCT 13160

Qy    580 TACTATAATGGTAATAATGGTAATACTCCTTTCAATCCAGGTAAAAGTTAATGCACACCT 639
          | |  ||   ||| || | ||| ||   ||     | | | |   | |  |||||||
Db  13161 TTCCCTACAAGTACTACTAGTACTAGCACTAGTGCTACTGTTCCTATTACAACCACACCT 13220
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Qy     640 TGTCCGGCAATTAAACCTGCTAATGTTGCTTAAGCTACTTTAGGTAATGATGCTACAATA 699
Db   13221 TCCCCTACAAATACTGCTGATGCTAACACTAGTAATACTGTTCCTAATACCACTATGCCT 13280

Qy     700 ACCGCATAATGTAACGTTGCATGCCCTGATGGTACTATAAGTGCTGCTGGAGTAAATAAT 759
Db   13281 TCTCCTACAAGTAGTACTACTGTGAGTACTATTGCTACCGTTCCCATTTCAGTGACTCCT 13340

Qy     760 TGGGTAGCACAAAACACTGAATGTACTAATTGTGCTCCTAACTTTTACAATAATAATGCT 819
Db   13341 TCTCTGACAAGTACTGCTGATGCCACCATTAGTACTACTGTACTTATTGCCACTACTTCT 13400

Qy     820 CCTAATTTCAATCCAGGTAATAGTACATGCCTACCTTGCCCAGCAAATAAAGATTATGGT 879
Db   13401 TCTCTAACAGGTACTACTGATGTTAGCACTAGTACTACTATTAATAATATAAGTACTCCT 13460

Qy     880 GCTGAAGCCACTGCAGGTGGTGCCGCTACTTTAGCCAAATAATGTAATATTGCATGCCCT 939
Db   13461 GTTCAAACAAATACTACTAATGCTAGCACTAGTACTAATGTTGCTAATATAACTGCTACC 13520

Qy     940 GATGGTACTGCAATTGCTAGTGGAGCAACTAATTATGTAATATTATAAACAGAATGTCTA 999
Db   13521 TCTCATACAAGTACTGATGATACTGTTCCTAATAATACTGTTCCAGTTACAGCTATTCCT 13580

Qy    1000 AATTGTGCTGCTA 1012
Db   13581 TCTCTTGCAAATA 13593
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RESULT 5
US-09-864-761-19241
; Sequence 19241, Application US/09864761
; Patent No. US20020048763A1
; GENERAL INFORMATION:
;  APPLICANT: Penn, Sharron G.
;  APPLICANT:  Rank, David R.
;  APPLICANT:  Hanzel, David K.
;  APPLICANT:  Chen, Wensheng
;  TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
;  TITLE OF INVENTION:  GENE EXPRESSION ANALYSIS BY MICROARRAY
;  FILE REFERENCE: Aeomica-X-1
;  CURRENT APPLICATION NUMBER: US/09/864,761
;  CURRENT FILING DATE:  2001-05-23
;  PRIOR APPLICATION NUMBER: US 60/180,312
;  PRIOR FILING DATE: 2000-02-04
;  PRIOR APPLICATION NUMBER: US 60/207,456
;  PRIOR FILING DATE: 2000-05-26
;  PRIOR APPLICATION NUMBER: US 09/632,366
;  PRIOR FILING DATE: 2000-08-03
;  PRIOR APPLICATION NUMBER: GB 24263.6
;  PRIOR FILING DATE: 2000-10-04
;  PRIOR APPLICATION NUMBER: US 60/236,359
;  PRIOR FILING DATE: 2000-09-27
;  PRIOR APPLICATION NUMBER: PCT/US01/00666
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00667
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00664
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00669
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00665
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00668
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00663
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00662
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00661
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00670
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: US 60/234,687
;  PRIOR FILING DATE: 2000-09-21
;  PRIOR APPLICATION NUMBER: US 09/608,408
;  PRIOR FILING DATE: 2000-06-30
;  PRIOR APPLICATION NUMBER: US 09/774,203
;  PRIOR FILING DATE: 2001-01-29
;  NUMBER OF SEQ ID NOS: 49117
;  SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
; SEQ ID NO 19241
;  LENGTH: 1075
;  TYPE: DNA
;  ORGANISM: Homo sapiens
;  FEATURE:
;  OTHER INFORMATION: MAP TO AL078472.1
;  OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 27
;  OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 19
;  OTHER INFORMATION: EXPRESSED IN BT474, SIGNAL = 34
;  OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 7.1
;  OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 28
;  OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 43
;  OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 20
;  OTHER INFORMATION: EXPRESSED IN HBL100, SIGNAL = 25
;  OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 18
;  OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 21
;  OTHER INFORMATION: EST_HUMAN HIT: AV739739.1, EVALUE 1.00e+00
;  OTHER INFORMATION: NT HIT: AL163201.2, EVALUE 2.00e-19
US-09-864-761-19241

  Query Match             3.3%;  Score 45.8;  DB 10;  Length 1075;
  Best Local Similarity   41.7%;  Pred. No. 0.098;
  Matches  287;  Conservative    0;  Mismatches  402;  Indels    0;  Gaps    0;

Qy     288 TGCTGGTACCGCAATTGCAGGTGGAGCAACAGATTATGCAGCAATAATCACAGAATGTGT 347
Db     216 TGATGGTGGTGATGGTGGTGGTGGTGATGGTGGTGGTGGTGGTGATGGTGATGGTGGTGG 275

Qy     348 TAATTGTAGAATTAATTTTTATAATGAAAATGCTCCAAATTTTAATGCAGGTGCTAGTAC 407
Db     276 TGGTGATGGTGGTGATGGTAGTGGTGATGGTGGTGGTGGTGGTGATGGTGGTGATGGTGA 335

Qy     408 ATGCACAGCTTGTCCGGTAAACAGAGTTGGTGGTGCATTGACTGCTGGTAATGCCGCTAC 467
Db     336 TGGTGGTGGTGGTGATGGTGGTGATGGTAGTGGTGGTAGTGGTACTGGTGGTGGTGGTGA 395

Qy     468 CATAGTCGCATAATGTAACGTCGCATGTCCTACTGGTACTGCACTTGATGATGGAGTAAC 527
Db     396 TGGTGGTGGTGCTGGTGATGGTGGTAGTGGTGGTGGTGGTAGTGGTGATGGTGGTGGTGG 455

Qy     528 TACTGATTATGTTAGATCATTCACAGAATGTGTTAAATGTAGACTTAACTTTTACTATAA 587
Db     456 TGGTGGTAGTGGTAGTGGTGATGGTGATGATGGTGGTGGTGGTGGTCGTGGTTTTAG 515

Qy     588 TGGTAATAATGGTAATACTCCTTTCAATCCAGGTAAAAGTTAATGCACACCTTGTCCGGC 647
Db     516 TGGTGATGGTGGTGGTGGTGGTGGTGGTAGTGGTGGTGATGCTGGTGGTGGTGGTGG 575

Qy     648 AATTAAACCTGCTAATGTTGCTTAAGCTACTTTAGGTAATGATGCTACAATAACCGCATA 707
Db     576 TAGTGGTGATGGTGGTGGTGGTGATGGTGGTGGTGGTGATGGTGGTGGTGGTGATGGTGG 635

Qy     708 ATGTAACGTTGCATGCCCTGATGGTACTATAAGTGCTGCTGGAGTAAATAATTGGGTAGC 767
Db     636 TGGTGGTGGTGGTAGTGGTGATGGTGGTGGTGGTGATGGTGGTGGTGGTGATGGTGGTGG 695

Qy     768 ACAAAACACTGAATGTACTAATTGTGCTCCTAACTTTTACAATAATAATGCTCCTAATTT 827
Db     696 TGGTGATGGTGGTGGTGGTGATGGTGGTGGTAGTGGTGGTAGTGGTGATGGTGGTAGTGG 755

Qy     828 CAATCCAGGTAATAGTACATGCCTACCTTGCCCAGCAAATAAAGATTATGGTGCTGAAGC 887
Db     756 TGATGGTGGTGGTGGTGATGGTGGTGGTGGTGGTGGTGGTAGTGGTGATGGTGGTAGTGG 815

Qy     888 CACTGCAGGTGGTGCCGCTACTTTAGCCAAATAATGTAATATTGCATGCCCTGATGGTAC 947
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Db    816 TCATGGTGGTGGTGGTGGTGGTCATGGTTTTAGTGGTGATGGTGGTGGTGGTGGTGGTAG 875

Qy    948 TGCAATTGCTAGTGGAGCAACTAATTATG 976
Db    876 TGGTGGTGGTGGTTGTGGTAGTGGTGGTG 904
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RESULT 6
US-09-864-761-2513
; Sequence 2513, Application US/09864761
; Patent No. US20020048763A1
; GENERAL INFORMATION:
; APPLICANT: Penn, Sharron G.
; APPLICANT: Rank, David R.
; APPLICANT: Hanzel, David K.
; APPLICANT: Chen, Wensheng
; TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
; TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
; FILE REFERENCE: Aeomica-X-1
; CURRENT APPLICATION NUMBER: US/09/864,761
; CURRENT FILING DATE: 2001-05-23
; PRIOR APPLICATION NUMBER: US 60/180,312
; PRIOR FILING DATE: 2000-02-04
; PRIOR APPLICATION NUMBER: US 60/207,456
; PRIOR FILING DATE: 2000-05-26
; PRIOR APPLICATION NUMBER: US 09/632,366
; PRIOR FILING DATE: 2000-08-03
; PRIOR APPLICATION NUMBER: GB 24263.6
; PRIOR FILING DATE: 2000-10-04
; PRIOR APPLICATION NUMBER: US 60/236,359
; PRIOR FILING DATE: 2000-09-27
; PRIOR APPLICATION NUMBER: PCT/US01/00666
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00667
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00664
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00669
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00665
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00668
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00663
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00662
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00661
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00670
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: US 60/234,687
; PRIOR FILING DATE: 2000-09-21
; PRIOR APPLICATION NUMBER: US 09/608,408
; PRIOR FILING DATE: 2000-06-30
; PRIOR APPLICATION NUMBER: US 09/774,203
; PRIOR FILING DATE: 2001-01-29
; NUMBER OF SEQ ID NOS: 49117
; SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
; SEQ ID NO 2513
; LENGTH: 1403
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; OTHER INFORMATION: MAP TO AL078472.1
; OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 27
; OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 19
; OTHER INFORMATION: EXPRESSED IN BT474, SIGNAL = 34
; OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 7.1
; OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 28
; OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 43
; OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 20
; OTHER INFORMATION: EXPRESSED IN HBL100, SIGNAL = 25
; OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 18
; OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 21
US-09-864-761-2513

Query Match 3.3%; Score 45.8; DB 10; Length 1403;
Best Local Similarity 41.7%; Pred. No. 0.11;
Matches 287; Conservative 0; Mismatches 402; Indels 0; Gaps 0;

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Qy    288 TGCTGGTACCGCAATTGCAGGTGGAGCAACAGATTATGCAGCAATAATCACAGAATGTGT 347
Db    566 TGATGGTGGTGATGGTGGTGGTGGTGATGGTGGTGGTGGTGGTGATGGTGATGGTGGTGG 625

Qy    348 TAATTGTAGAATTAATTTTTATAATGAAAATGCTCCAAATTTTAATGCAGGTGCTAGTAC 407
Db    626 TGGTGATGGTGGTGATGGTAGTGGTGATGGTGGTGGTGGTGGTGATGGTGGTGATGGTGA 685

Qy    408 ATGCACAGCTTGTCCGGTAAACAGAGTTGGTGGTGCATTGACTGCTGGTAATGCCGCTAC 467
Db    686 TGGTGGTGGTGGTGATGGTGGTGATGGTAGTGGTGGTAGTGGTACTGGTGGTGGTGGTGA 745

Qy    468 CATAGTCGCATAATGTAACGTCGCATGTCCTACTGGTACTGCACTTGATGATGGAGTAAC 527
Db    746 TGGTGGTGGTGCTGGTGATGGTGGTAGTGGTGGTGGTGGTAGTGGTGATGGTGGTGGTGG 805

Qy    528 TACTGATTATGTTAGATCATTCACAGAATGTGTTAAATGTAGACTTAACTTTTACTATAA 587
Db    806 TGGTGGTAGTGGTAGTGGTGATGGTGATGATGGTGGTGGTGGTGGTGGTCGTGGTTTTAG 865

Qy    588 TGGTAATAATGGTAATACTCCTTTCAATCCAGGTAAAAGTTAATGCACACCTTGTCCGGC 647
Db    866 TGGTGATGGTGGTGGTGGTGGTGGTGGTGGTAGTGGTGGTGATGCTGGTGGTGGTGGTGG 925

Qy    648 AATTAAACCTGCTAATGTTGCTTAAGCTACTTTAGGTAATGATGCTACAATAACCGCATA 707
Db    926 TAGTGGTGATGGTGGTGGTGGTGATGGTGGTGGTGGTGATGGTGGTGGTGGTGATGGTGG 985

Qy    708 ATGTAACGTTGCATGCCCTGATGGTACTATAAGTGCTGCTGGAGTAAATAATTGGGTAGC 767
Db    986 TGGTGGTGGTGGTAGTGGTGATGGTGGTGGTGGTGATGGTGGTGGTGGTGATGGTGGTGG 1045

Qy    768 ACAAAACACTGAATGTACTAATTGTGCTCCTAACTTTTACAATAATAATGCTCCTAATTT 827
Db   1046 TGGTGATGGTGGTGGTGGTGATGGTGGTGGTAGTGGTGGTAGTGGTGATGGTGGTAGTGG 1105

Qy    828 CAATCCAGGTAATAGTACATGCCTACCTTGCCCAGCAAATAAAGATTATGGTGCTGAAGC 887
Db   1106 TGATGGTGGTGGTGGTGATGGTGGTGGTGGTGGTGGTGGTAGTGGTGATGGTGGTAGTGG 1165

Qy    888 CACTGCAGGTGGTGCCGCTACTTTAGCCAAATAATGTAATATTGCATGCCCTGATGGTAC 947
Db   1166 TCATGGTGGTGGTGGTGGTGGTCATGGTTTTAGTGGTGATGGTGGTGGTGGTGGTGGTAG 1225

Qy    948 TGCAATTGCTAGTGGAGCAACTAATTATG 976
Db   1226 TGGTGGTGGTGGTTGTGGTAGTGGTGGTG 1254
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RESULT 7
US-09-798-042-95/c
; Sequence 95, Application US/09798042
; Patent No. US20020068343A1
; GENERAL INFORMATION:
; APPLICANT: Reed, Steven G.
; APPLICANT: Lodes, Michael J.
; APPLICANT: Houghton, Raymond L.
; APPLICANT: McNeill, Patricia D.
; TITLE OF INVENTION: COMPOUNDS AND METHODS FOR THE DIAGNOSIS
; TITLE OF INVENTION: AND TREATMENT OF EHRLICHIA INFECTION
; FILE REFERENCE: 210121.439C7
; CURRENT APPLICATION NUMBER: US/09/798,042
; CURRENT FILING DATE: 2001-03-02
; NUMBER OF SEQ ID NOS: 108

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;  SOFTWARE:  FastSEQ for Windows Version 3.0
; SEQ ID NO 95
;   LENGTH: 2120
;   TYPE: DNA
;   ORGANISM: Ehrlichia
US-09-798-042-95

  Query Match             3.1%;  Score 43.4;  DB 10;  Length 2120;
  Best Local Similarity   44.3%;  Pred. No. 0.5;
  Matches  277;  Conservative    0;  Mismatches  336;  Indels   12;  Gaps    2;

Qy     287 CTGCTGGTACCGCAATTGCAGGTGGAGCAACAGATTATGCAGCAATAATCACAGAATGTG 346
           || || |   | |  |||   |  | |||||| | |||  | |  ||  | | |
Db    1502 CTTCTTGCTGCATACCAGCATCAGCTTCTACAGATACTTCAGACTTCAGATCACCTTCAG 1443

Qy     347 TTAATTGTAGAATTAATTTTTATAATGAAAATGCTCCAAATTTTAATGCAGGTGCTAGTA 406
           | |     |||| |     |    |     || | || |   |   | || || |  ||
Db    1442 TAACACCAAGAACTGTAGACTCAGGTTGTGCTGGTGCAGAAACTTCAGGAGCTGATTCTA 1383

Qy     407 CATGCACAGCTTGTCCGGTAAACAGAGTTGGTGGTGCATTGACTGCTGGTAATGCCGCTA 466
             ||    |||| |   | ||  |    | | |    ||   | ||  || ||  | ||
Db    1382 GTTGTTGCGCTTCTGGAGCAACTACCACTTCTTGAAGCTTATTTTCTTCTAGTGATGGTA 1323

Qy     467 CCATAGTCGCATAATGTAACGTCGCATGTCCTACTGGTACTGCACTTGATGATGGAGTAA 526
           | || |   |   | |       || ||  | ||| | | ||   |   | |
Db    1322 CAATCGCTTCTGCAGCTTCAACACCAGGTAATTCTGCTTCAGCTACTACAGGTACTTGTG 1263

Qy     527 CTACTGATTATGTTAGATCATTCACAGAATGT---GTTAAATGTAGACTTAACTTTTACT 583
           |||| | ||     ||||| ||| || || |    ||| | | |    |   | ||  |
Db    1262 CTACAGGTTGCTCAAGATCTATCAAAGTATCTTCCTTTAGAAGAACTTCTGTTTCTTCTT 1203

Qy     584 ATAATGGTAATAATGGTAATACTCCTTTCAATCCAGGTAAAAGTTAATGCACACCTTGTC 643
             || |  ||     | |     ||| |  | | |   |  || ||  ||| |   ||| |
Db    1202 TTACTTCTACAGGAGCTTCAGTTCCCTCTAGTGCTTCTGCAATTTCTTGCTCTTGTTGAC 1143

Qy     644 CGGCAATTAAACCTGCTAATGTTGCTTAAGCTACTTTAGGTAATGATGCTACAATAACCG 703
           | |  ||||   ||  |   | | | | |||          ||  ||| || ||
Db    1142 CAGAGATTACTTCTTTTTGCGCTACATCAGCATTAGCTTCTACAGATACTTCAGACTTTA 1083

Qy     704 CATAATGTAACGTTGCATGCCCTGATGGTACTATAAGTGCTGCTGGAGTAAATAATTGGG 763
           || |  |   |  ||         ||          | ||  |||||| | | | | ||  |
Db    1082 GATCACCTTCAGCAACACCAAGAACTGTAGACTCAGGTTGTGCTGGCGCAGAAACTTCAG 1023

Qy     764 TAGCACAAAACACTGAATGTACTAATTGTGCTCCTAACTTTTACAATAATAATGCTCCTA 823
             ||          |||| | || |  ||| ||| ||          | | |  |
Db    1022 GAG---------CTGATTCTAGTTGTTGCGCTTCTGGAGCAACTACCACTTCTTGAAGCT 972

Qy     824 ATTTCAATCCAGGTAATAGTACATGCCTACCTTGCCCAGCAAATAAAGATTATGGTGCTG 883
             ||   | |  || || ||||| |    ||    |  |||    || | ||  ||||
Db     971 TATTTTCTTCTAGTGATGGTACAATCGCTTCTGCAGCTTCAACACCAGGTAATTCTGCTT 912

Qy     884 AAGCCACTGCAGGTGGTGCCGCTAC 908
           |||  ||| |||||  |  ||||||
Db     911 CAGCTACTACAGGTACTTGCGCTAC 887


RESULT 8
US-09-159-469-39
; Sequence 39, Application US/09159469
; Patent No. US20020064535A1
;  GENERAL INFORMATION:
;    APPLICANT:  Reed, Steven G.
;    APPLICANT:  Lodes, Michael J.
;    APPLICANT:  Houghton, Raymond
;    TITLE OF INVENTION:  COMPOUNDS AND METHODS FOR THE DIAGNOSIS AND
;    TITLE OF INVENTION:  THERAPY OF EHRLICHIA INFECTION
;    NUMBER OF SEQUENCES:  73
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  SEED and BERRY LLP
;      STREET:  6300 Columbia Center, 701 Fifth Avenue
;      CITY:  Seattle
;      STATE:  Washington
;      COUNTRY:  USA
;      ZIP:  98104
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/159,469
;      FILING DATE:
;      CLASSIFICATION:
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  09/106,582
;      FILING DATE:  29-JUN-1998
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Maki, David J.
;      REGISTRATION NUMBER:  31,392
;      REFERENCE/DOCKET NUMBER:  210121.439C2
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  206-622-4900
;      TELEFAX:  206-682-6031
;  INFORMATION FOR SEQ ID NO:  39:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  2129 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  DNA (genomic)
US-09-159-469-39

  Query Match             3.1%;  Score 43.4;  DB 10;  Length 2129;
  Best Local Similarity   44.3%;  Pred. No. 0.5;
  Matches  277;  Conservative    0;  Mismatches  336;  Indels   12;  Gaps    2;

Qy     287 CTGCTGGTACCGCAATTGCAGGTGGAGCAACAGATTATGCAGCAATAATCACAGAATGTG 346
           || || |   | |  |||   |  | |||||| | |||  | |  ||  | | |
Db     619 CTTCTTGCTGCATACCAGCATCAGCTTCTACAGATACTTCAGACTTCAGATCACCTTCAG 678

Qy     347 TTAATTGTAGAATTAATTTTTATAATGAAAATGCTCCAAATTTTAATGCAGGTGCTAGTA 406
           | |     |||| |     |    |     || | || |   |   | || || |  ||
Db     679 TAACACCAAGAACTGTAGACTCAGGTTGTGCTGGTGCAGAAACTTCAGGAGCTGATTCTA 738

Qy     407 CATGCACAGCTTGTCCGGTAAACAGAGTTGGTGGTGCATTGACTGCTGGTAATGCCGCTA 466
             ||    |||| |   | ||  |    | | |    ||   | ||  || ||  | ||
Db     739 GTTGTTGCGCTTCTGGAGCAACTACCACTTCTTGAAGCTTATTTTCTTCTAGTGATGGTA 798

Qy     467 CCATAGTCGCATAATGTAACGTCGCATGTCCTACTGGTACTGCACTTGATGATGGAGTAA 526
           | || |   |   | |       || ||  | ||| | | ||   |   | |
Db     799 CAATCGCTTCTGCAGCTTCAACACCAGGTAATTCTGCTTCAGCTACTACAGGTACTTGTG 858

Qy     527 CTACTGATTATGTTAGATCATTCACAGAATGT---GTTAAATGTAGACTTAACTTTTACT 583
           |||| | ||     ||||| ||| || || |    ||| | | |    |   | ||  |
Db     859 CTACAGGTTGCTCAAGATCTATCAAAGTATCTTCCTTTAGAAGAACTTCTGTTTCTTCTT 918

Qy     584 ATAATGGTAATAATGGTAATACTCCTTTCAATCCAGGTAAAAGTTAATGCACACCTTGTC 643
             || |  ||     | |     ||| |  | | |   |  || ||  ||| |   ||| |
Db     919 TTACTTCTACAGGAGCTTCAGTTCCCTCTAGTGCTTCTGCAATTTCTTGCTCTTGTTGAC 978

Qy     644 CGGCAATTAAACCTGCTAATGTTGCTTAAGCTACTTTAGGTAATGATGCTACAATAACCG 703
           | |  ||||   ||  |   | | | | |||          ||  ||| || ||
Db     979 CAGAGATTACTTCTTTTTGCGCTACATCAGCATTAGCTTCTACAGATACTTCAGACTTTA 1038

Qy     704 CATAATGTAACGTTGCATGCCCTGATGGTACTATAAGTGCTGCTGGAGTAAATAATTGGG 763
           || |  |   |  ||         ||          | ||  |||||| | | | | ||  |
Db    1039 GATCACCTTCAGCAACACCAAGAACTGTAGACTCAGGTTGTGCTGGCGCAGAAACTTCAG 1098

Qy     764 TAGCACAAAACACTGAATGTACTAATTGTGCTCCTAACTTTTACAATAATAATGCTCCTA 823
             ||          |||| | || |  ||| ||| ||          | | |  |
Db    1099 GAG---------CTGATTCTAGTTGTTGCGCTTCTGGAGCAACTACCACTTCTTGAAGCT 1149

Qy     824 ATTTCAATCCAGGTAATAGTACATGCCTACCTTGCCCAGCAAATAAAGATTATGGTGCTG 883
             ||   | |  || || ||||| |    ||    |  |||    || | ||  ||||
Db    1150 TATTTTCTTCTAGTGATGGTACAATCGCTTCTGCAGCTTCAACACCAGGTAATTCTGCTT 1209
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Qy     884 AAGCCACTGCAGGTGGTGCCGCTAC 908
           ||| ||| |||||  |  ||||||
Db    1210 CAGCTACTACAGGTACTTGCGCTAC 1234
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RESULT 9
US-09-798-042-39
; Sequence 39, Application US/09798042
; Patent No. US20020068343A1
; GENERAL INFORMATION:
; APPLICANT: Reed, Steven G.
; APPLICANT: Lodes, Michael J.
; APPLICANT: Houghton, Raymond L.
; APPLICANT: McNeill, Patricia D.
; TITLE OF INVENTION: COMPOUNDS AND METHODS FOR THE DIAGNOSIS
; TITLE OF INVENTION: AND TREATMENT OF EHRLICHIA INFECTION
; FILE REFERENCE: 210121.439C7
; CURRENT APPLICATION NUMBER: US/09/798,042
; CURRENT FILING DATE: 2001-03-02
; NUMBER OF SEQ ID NOS: 108
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 39
; LENGTH: 2129
; TYPE: DNA
; ORGANISM: Ehrlichia sp.
US-09-798-042-39

Query Match 3.1%; Score 43.4; DB 10; Length 2129;
Best Local Similarity 44.3%; Pred. No. 0.5;
Matches 277; Conservative 0; Mismatches 336; Indels 12; Gaps 2;

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Qy     287 CTGCTGGTACCGCAATTGCAGGTGGAGCAACAGATTATGCAGCAATAATCACAGAATGTG 346
           || || |   |  |   |||   |   | ||||||  | |||   | |   ||   |  |
Db     619 CTTCTTGCTGCATACCAGCATCAGCTTCTACAGATACTTCAGACTTCAGATCACCTTCAG 678

Qy     347 TTAATTGTAGAATTAATTTTTATAATGAAAATGCTCCAAATTTTAATGCAGGTGCTAGTA 406
           | |     |||| |      |    |     || | || |   |   | || || | ||
Db     679 TAACACCAAGAACTGTAGACTCAGGTTGTGCTGGTGCAGAAACTTCAGGAGCTGATTCTA 738

Qy     407 CATGCACAGCTTGTCCGGTAAACAGAGTTGGTGGTGCATTGACTGCTGGTAATGCCGCTA 466
           ||    |||| |   | || |    |  | |    ||   | ||  || ||   | ||
Db     739 GTTGTTGCGCTTCTGGAGCAACTACCACTTCTTGAAGCTTATTTTCTTCTAGTGATGGTA 798

Qy     467 CCATAGTCGCATAATGTAACGTCGCATGTCCTACTGGTACTGCACTTGATGATGGAGTAA 526
           | || |   |   |  |       || ||  | ||| | | ||   |   | |
Db     799 CAATCGCTTCTGCAGCTTCAACACCAGGTAATTCTGCTTCAGCTACTACAGGTACTTGTG 858

Qy     527 CTACTGATTATGTTAGATCATTCACAGAATGT---GTTAAATGTAGACTTAACTTTTACT 583
           |||| | ||     |||||  ||| || || |    ||| | | |    |   | ||  |
Db     859 CTACAGGTTGCTCAAGATCTATCAAAGTATCTTCCTTTAGAAGAACTTCTGTTTCTTCTT 918

Qy     584 ATAATGGTAATAATGGTAATACTCCTTTCAATCCAGGTAAAAGTTAATGCACACCTTGTC 643
           || |  ||     | |     ||| |  | | |   |  || ||  ||| |   ||| |
Db     919 TTACTTCTACAGGAGCTTCAGTTCCCTCTAGTGCTTCTGCAATTTCTTGCTCTTGTTGAC 978

Qy     644 CGGCAATTAAACCTGCTAATGTTGCTTAAGCTACTTTAGGTAATGATGCTACAATAACCG 703
           | |  ||||   ||  |   | | | | |||         ||  ||| || ||
Db     979 CAGAGATTACTTCTTTTTGCGCTACATCAGCATTAGCTTCTACAGATACTTCAGACTTTA 1038

Qy     704 CATAATGTAACGTTGCATGCCCTGATGGTACTATAAGTGCTGCTGGAGTAAATAATTGGG 763
           || |  |   |  ||        ||       | ||  |||||| | | | | ||  |
Db    1039 GATCACCTTCAGCAACACCAAGAACTGTAGACTCAGGTTGTGCTGGCGCAGAAACTTCAG 1098

Qy     764 TAGCACAAAACACTGAATGTACTAATTGTGCTCCTAACTTTTACAATAATAATGCTCCTA 823
           ||         |||| | || |  ||| ||| ||        |  | |  |
Db    1099 GAG---------CTGATTCTAGTTGTTGCGCTTCTGGAGCAACTACCACTTCTTGAAGCT 1149

Qy     824 ATTTCAATCCAGGTAATAGTACATGCCTACCTTGCCCAGCAAATAAAGATTATGGTGCTG 883
           ||   | |  || || |||||  |    ||    |  |||    || | ||  ||||
Db    1150 TATTTTCTTCTAGTGATGGTACAATCGCTTCTGCAGCTTCAACACCAGGTAATTCTGCTT 1209

Qy     884 AAGCCACTGCAGGTGGTGCCGCTAC 908
           ||| ||| |||||  |  ||||||
Db    1210 CAGCTACTACAGGTACTTGCGCTAC 1234
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RESULT 10
US-09-864-761-20174/c
; Sequence 20174, Application US/09864761
; Patent No. US20020048763A1
; GENERAL INFORMATION:
; APPLICANT: Penn, Sharron G.
; APPLICANT: Rank, David R.
; APPLICANT: Hanzel, David K.
; APPLICANT: Chen, Wensheng
; TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FO
; TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
; FILE REFERENCE: Aeomica-X-1
; CURRENT APPLICATION NUMBER: US/09/864,761
; CURRENT FILING DATE: 2001-05-23
; PRIOR APPLICATION NUMBER: US 60/180,312
; PRIOR FILING DATE: 2000-02-04
; PRIOR APPLICATION NUMBER: US 60/207,456
; PRIOR FILING DATE: 2000-05-26
; PRIOR APPLICATION NUMBER: US 09/632,366
; PRIOR FILING DATE: 2000-08-03
; PRIOR APPLICATION NUMBER: GB 24263.6
; PRIOR FILING DATE: 2000-10-04
; PRIOR APPLICATION NUMBER: US 60/236,359
; PRIOR FILING DATE: 2000-09-27
; PRIOR APPLICATION NUMBER: PCT/US01/00666
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00667
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00664
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00669
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00665
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00668
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00663
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00662
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00661
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00670
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: US 60/234,687
; PRIOR FILING DATE: 2000-09-21
; PRIOR APPLICATION NUMBER: US 09/608,408
; PRIOR FILING DATE: 2000-06-30
; PRIOR APPLICATION NUMBER: US 09/774,203
; PRIOR FILING DATE: 2001-01-29
; NUMBER OF SEQ ID NOS: 49117
; SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
; SEQ ID NO 20174
; LENGTH: 439
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; OTHER INFORMATION: MAP TO AL035419.9
; OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 1.3
; OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 1.2
; OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 1
; OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 1.4
; OTHER INFORMATION: EXPRESSED IN HBL100, SIGNAL = 1.1
; OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 1.5
; OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 1.6
; OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 1.5
; OTHER INFORMATION: EXPRESSED IN BT474, SIGNAL = 1.5
; OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 1.2
US-09-864-761-20174

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  Query Match             3.0%;  Score 42.8;  DB 10;  Length 439;
  Best Local Similarity   45.0%;  Pred. No. 0.35;
  Matches  161;  Conservative    0;  Mismatches  197;  Indels    0;  Gaps    0;

Qy     726 TGATGGTACTATAAGTGCTGCTGGAGTAAATAATTGGGTAGCACAAAACACTGAATGTAC 785
           || ||||  |     ||||||||  |   || || | |  |   | |  |||   | |
Db     382 TGGTGGTGATGGTGATGCTGCTGCTGATGATGATGGTGGGGATAATGATGGTGATGCTGC 323

Qy     786 TAATTGTGCTCCTAACTTTTACAATAATAATGCTCCTAATTTCAATCCAGGTAATAGTAC 845
             || ||| |  |      | |  ||  | ||||| || ||     |  ||  ||  | |
Db     322 CGATGGTGATGGTGGTGATGATGATGGTGATGCTGCTGATGGTGTTGATGGCGATGCTGC 263

Qy     846 ATGCCTACCTTGCCCAGCAAATAAAGATTATGGTGCTGAAGCCACTGCAGGTGGTGCCGC 905
                 |     |   || | |||  || |||||  ||  |||| ||||||||   |
Db     262 TGATGGTGATGATGGTGGTGATTACGATGGTGATGCTGCTGCTGCTGCTGGTGGTGATCC 203

Qy     906 TACTTTAGCCAAATAATGTAATATTGCATGCCCTGATGGTACTGCAATTGCTAGTGGAGC 965
           | ||   |       |  || ||  ||   |   || || |   |    || | |||  |
Db     202 TGCTGATGGTTTTGATGGTTATGATGATGGTGATGCTGCTGACGGTGATGATGGTGATGA 143

Qy     966 AACTAATTATGTAATATTATAAACAGAATGTCTAAATTGTGCTGCTAACTTTTATTTTGA 1025
             | ||  |          |    ||  ||   ||| ||  || |      |     |||
Db     142 TGATGATGGTAATGATGATGATGGTGAGGGTAATAATGGTAATGATGGTGATGGCGATGA 83

Qy    1026 TGGTAATAATTTCTAGGCAGGAAGTAGTAGATGCAAAGCATGTCCAGCAAATAAAGTT 1083
           |||| |||||  | | |   |    | ||      || |   ||   || | |   |||
Db      82 TGGTGATAATAACGATGATGCTGATGGTGATGATAATGATGGTGATGCTACTGCTGTT 25


RESULT 11
US-09-970-477-1
; Sequence 1, Application US/09970477
; Patent No. US20020127545A1
; GENERAL INFORMATION:
;  APPLICANT: Lorincz, Attila T.
;  TITLE OF INVENTION: ASSESSMENTS OF HUMAN PAPILLOMA VIRUS-RELATED DISEASE
;  FILE REFERENCE: 2629-4005US4
;  CURRENT APPLICATION NUMBER: US/09/970,477
;  CURRENT FILING DATE:  2001-10-04
;  PRIOR APPLICATION NUMBER: U.S. 09/210,168
;  PRIOR FILING DATE: 1998-12-11
;  PRIOR APPLICATION NUMBER: U.S. 60/082,167
;  PRIOR FILING DATE: 1998-04-17
;  PRIOR APPLICATION NUMBER: U.S. 60/070,486
;  PRIOR FILING DATE: 1998-01-05
;  PRIOR APPLICATION NUMBER: U.S. 60/069,426
;  PRIOR FILING DATE: 1997-12-12
;  NUMBER OF SEQ ID NOS: 5
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
;   LENGTH: 1390
;   TYPE: DNA
;   ORGANISM: Human papillomavirus
;   FEATURE:
;   OTHER INFORMATION: L2-HPV16
US-09-970-477-1

  Query Match             3.0%;  Score 42.6;  DB 10;  Length 1390;
  Best Local Similarity   44.1%;  Pred. No. 0.65;
  Matches  226;  Conservative    0;  Mismatches  284;  Indels    3;  Gaps    1;

Qy     633 CACACCTTGTCCGGCAATTAAACCTGCTAATGTTGCTTAAGCTACTTTAGGTAATGATGC 692
           ||| ||   |      |||| |  || ||||  ||| || |    | ||| |   |||
Db     834 CACTCCCACTAAACTTATTACATATGATAATCCTGCATATGAAGGTATAGATGTGGATAA 893

Qy     693 TACAATAACCGCATAATGTAACGTTGCATGCCCTGATGGTACTATAAGTGCTGCTGGAGT 752
           |||| ||      |   |||| | |    |  | ||   ||  |  | |||      |
Db     894 TACATTATATTTTTCTAGTAATGATAATAGTATTAATATAGCTCCAGATCCTGACTTTTT 953

Qy     753 AAATAATTGGGTAGCACAAAACACTGAATGTACTAATTGTGCTCCTAACTTT---TACAA 809
             |||     |    ||| |   | | ||  ||   | |   | ||  | ||   ||||
Db     954 GGATATAGTTGCTTTACATAGGCCAGCATTAACCTCTAGGCGTACTGGCATTAGGTACAG 1013

Qy     810 TAATAATGCTCCTAATTTCAATCCAGGTAATAGTACATGCCTACCTTGCCCAGCAAATAA 869
           ||  | || |  |||     |  | | ||| | |||   |   | ||       || ||
Db    1014 TAGAATTGGTAATAAACAAACACTACGTACTCGTAGTGGAAAATCTATAGGTGCTAAGGT 1073

Qy     870 AGATTATGGTGCTGAAGCCACTGCAGGTGGTGCCGCTACTTTAGCCAAATAATGTAATAT 929
           | |||||  |  |||     | | |   || | | ||       |    ||| |   | |||
Db    1074 ACATTATTATTATGATTTAAGTACTATTGATCCTGCAGAAGAAATAGAATTACAAACTAT 1133

Qy     930 TGCATGCCCTGATGGTACTGCAATTGCTAGTGGAGCAACTAATTATGTAATATTATAAAC 989
             ||    ||      |||| | | |   || |||  |   |  |   ||     |
Db    1134 AACACCTTCTACATATACTACCACTTCACATGCAGCCTCACCTACTTCTATTAATAATGG 1193

Qy     990 AGAATGTCTAAATTGTGCTGCTAACTTTTATTTTGATGGTAATAATTTCTAGGCAGGAAG 1049
           |  || |   | || ||| | | |||||  |   |||  |  ||    |  || |  |
Db    1194 ATTATATGATATTTATGCAGATGACTTTATTACAGATACTTCTACAACCCCGGTACCATC 1253

Qy    1050 TAGTAGATGCAAAGCATGTCCAGCAAATAAAGTTTAAGGCGCTGTAGCAACTGCAGGTGG 1109
           |      | | | | |   |||  |||    |  |    |   |   ||  |||||
Db    1254 TGTACCCTCTACATCTTTATCAGGTTATATTCCTGCAAATACAACAATTCCTTTTGGTGG 1313

Qy    1110 TACTGCTACTTTAATTGCATAATGTGCCCTTGA 1142
           | |   | | |   |  |  ||  |  | |||
Db    1314 TGCATACAATATTCCTTTAGTATCAGGTCCTGA 1346


RESULT 12
US-10-024-623-31/c
; Sequence 31, Application US/10024623
; Publication No. US20020187524A1
; GENERAL INFORMATION:
;  APPLICANT: Curtis, Rory A.J.
;  TITLE OF INVENTION: 8099, 46455, 54414, 53763, 67076, 67102, 44181,
;  TITLE OF INVENTION:  67084FL, AND 67084 ALT, HUMAN PROTEINS AND METHODS OF
;  TITLE OF INVENTION:  USE THEREOF
;  FILE REFERENCE: MNI-214CP
;  CURRENT APPLICATION NUMBER: US/10/024,623
;  CURRENT FILING DATE:  2001-12-17
;  PRIOR APPLICATION NUMBER: US 60/256,240
;  PRIOR FILING DATE: 2000-12-15
;  PRIOR APPLICATION NUMBER: US 60/256,588
;  PRIOR FILING DATE: 2000-12-18
;  PRIOR APPLICATION NUMBER: US 60/258,028
;  PRIOR FILING DATE: 2000-12-21
;  NUMBER OF SEQ ID NOS: 40
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 31
;   LENGTH: 25002
;   TYPE: DNA
;   ORGANISM: Caenorhabditis elegans
US-10-024-623-31

  Query Match             3.0%;  Score 42.6;  DB 9;  Length 25002;
  Best Local Similarity   52.5%;  Pred. No. 2.3;
  Matches  116;  Conservative    0;  Mismatches  104;  Indels    1;  Gaps    1;

Qy       1 ATGAAAAATAATATTTTAGTAATATTGATTATTTCATTATTTATCAATTAAATTAAATCT 60
           ||| | | |   |  ||| |  |||| || | |  |||  ||  | || | |||| || |
Db    4751 ATGTACATTTCAAAATTATTTTTATTCATCAATCTATTTATTCACTATCATATTAGAT-T 4693

Qy      61 GCTAATTGTCCTGTTGGAACTGAAACTAACACAGCCGGATAAGTTGATGATCTAGGAACT 120
           ||  || | |      | ||||   |||   |  |  |||  ||  |  | ||  | ||
Db    4692 TCTTTTTCTAATCCATAATCTGATCATAAATTACACTAATACATTTCTTCTTTAAAACCT 4633

Qy     121 CCTGCAAATTGTGTTAATTGTTAGAAAAACTTTTATTATAATAATGCTGCTGCTTTCGTT 180
           |  |     |||||||  |   ||   | |  |||  | ||  |||  |  | |||| |||
Db    4632 CAGGATCCTTGTGTTTTTGACTATGTATAAGTTTGATGTATAAATAGTAATCCTTTTGTT 4573

Qy     181 CCTGGTGCTAGTACGTGTACACCTTGTCCATAAAAAAAAGA 221
           |||  || |  |       |||| |||   ||  | || | |
Db    4572 CCTTCTGTTGATTTTCCAACACATTGAATATTGATAATAAA 4532


RESULT 13
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US-09-864-761-228/c
; Sequence 228, Application US/09864761
; Patent No. US20020048763A1
; GENERAL INFORMATION:
;  APPLICANT: Penn, Sharron G.
;  APPLICANT:  Rank, David R.
;  APPLICANT:  Hanzel, David K.
;  APPLICANT:  Chen, Wensheng
;  TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
;  TITLE OF INVENTION:  GENE EXPRESSION ANALYSIS BY MICROARRAY
;  FILE REFERENCE: Aeomica-X-1
;  CURRENT APPLICATION NUMBER: US/09/864,761
;  CURRENT FILING DATE:  2001-05-23
;  PRIOR APPLICATION NUMBER: US 60/180,312
;  PRIOR FILING DATE: 2000-02-04
;  PRIOR APPLICATION NUMBER: US 60/207,456
;  PRIOR FILING DATE: 2000-05-26
;  PRIOR APPLICATION NUMBER: US 09/632,366
;  PRIOR FILING DATE: 2000-08-03
;  PRIOR APPLICATION NUMBER: GB 24263.6
;  PRIOR FILING DATE: 2000-10-04
;  PRIOR APPLICATION NUMBER: US 60/236,359
;  PRIOR FILING DATE: 2000-09-27
;  PRIOR APPLICATION NUMBER: PCT/US01/00666
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00667
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00664
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00669
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00665
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00668
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00663
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00662
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00661
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00670
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: US 60/234,687
;  PRIOR FILING DATE: 2000-09-21
;  PRIOR APPLICATION NUMBER: US 09/608,408
;  PRIOR FILING DATE: 2000-06-30
;  PRIOR APPLICATION NUMBER: US 09/774,203
;  PRIOR FILING DATE: 2001-01-29
;  NUMBER OF SEQ ID NOS: 49117
;  SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
; SEQ ID NO 228
;   LENGTH: 574
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: MAP TO AL079301.14
;   OTHER INFORMATION: EXPRESSED IN HBL100, SIGNAL = 7.8
;   OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 5.2
;   OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 8.2
;   OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 8.4
;   OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 4.7
;   OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 4.3
;   OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 5.6
;   OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 6.2
;   OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 7.8
;   OTHER INFORMATION: EXPRESSED IN BT474, SIGNAL =   7
US-09-864-761-228

  Query Match             3.0%;  Score 42;  DB 10;  Length 574;
  Best Local Similarity   42.7%;  Pred. No. 0.61;
  Matches  216;  Conservative    0;  Mismatches  290;  Indels    0;  Gaps    0;

Qy    345 TGTTAATTGTAGAATTAATTTTTATAATGAAAATGCTCCAAATTTTAATGCAGGTGCTAG 404
          || | || || |   | ||  || | || |  ||| |   |||  | ||   ||| | |
Db    532 TGGTGATGGTGGTGATGATGGTTGTCATAATGATGGTGATAATGGTGATCATTGTGATGG 473

Qy    405 TACATGCACAGCTTGTCCGGTAAACAGAGTTGGTGGTGCATTGACTGCTGGTAATGCCGC 464
          ||   ||   | |  | | |   |  | | || |  ||        || || | |||  |
Db    472 TAGTGGCGGTGATGCTGCTGGTGATGGGGGTGCTCATGGTGATGGTGGTGCTCATGGTGT 413

Qy    465 TACCATAGTCGCATAATGTAACGTCGCATGTCCTACTGGTACTGCACTTGATGATGGAGT 524
          |      |  ||      || | |        |  |  ||||  ||     |||| | |
Db    412 TGGTGGTGCTGCTGCTGGTGATGGTTGTCATGGTGATGGTGGTGATGAGGATGGTTGTCA 353

Qy    525 AACTACTGATTATGTTAGATCATTCACAGAATGTGTTAAATGTAGACTTAACTTTTACTA 584
          | |  || | ||  | |        ||  ||  |    || |   | |   | |
Db    352 TAGTGATGGTGATAATGGTGATCATTGTGATGGTAGTGGTGGTGGTGCTGATGGTGATGG 293

Qy    585 TAATGGTAATAATGGTAATACTCCTTTCAATCCAGGTAAAAGTTAATGCACACCTTGTCC 644
          |  || |  |  |||| ||  |  |     |  ||| |     |  |    | | ||
Db    292 TGGTGCTGCTGCTGGTGATGGTTGTCATGGTGATGGTGATGAGGATGGTTGTCATAGTGT 233

Qy    645 GGCAATTAAACCTGCTAATGTTGCTTAAGCTACTTTAGGTAATGATGCTACAATAACCGC 704
          |   |||  || | ||| || |     |  |  |||  |||||  |          ||
Db    232 TGGTGATAATGGTGATCATGGTGATGGTAGTGGTGGTGGTGCTGATGGTGATGGTGGTGC 173

Qy    705 ATAATGTAACGTTGCATGCCCTGATGGTACTATAAGTGCTGCTGGAGTAAATAATTGGGT 764
          |  || | | ||   |  ||| ||||  |    ||| || ||| |   ||  |   |
Db    172 TCATGGTGATGGTGGTGGTGCTGCTGGTGATGGTGGTGGTGATGGTGATCATTGTGATGG 113

Qy    765 AGCACAAAACACTGAATGTACTAATTGTGCTCCTAACTTTTACAATAATAATGCTCCTAA 824
                  ||    |  | || ||| |  | |   |    |||   | | |  |||
Db    112 TAGTGGCGGTGGTGCCCATGGTGATGGTGATGGTGATGGTGTTGATAGCTACGGTGATAA 53

Qy    825 TTTCAATCCAGGTAATAGTACATGCC 850
          |||  |   ||| || | |  || |
Db     52 CATCAGTGGTGGTCATGGCAGCTGTC 27


RESULT 14
US-09-938-842A-3346
; Sequence 3346, Application US/09938842A
; Patent No. US20020160378A1
; GENERAL INFORMATION:
;  APPLICANT: Harper, Jeff
;  APPLICANT:  Kreps, Joel
;  APPLICANT:  Wang, Xun
;  APPLICANT:  Zhu, Tong
;  TITLE OF INVENTION: STRESS-REGULATED GENES OF PLANTS, TRANSGENIC PLANTS CONTAINING
;  TITLE OF INVENTION:  SAME, AND METHODS OF USE
;  FILE REFERENCE: SCRIP1300-3
;  CURRENT APPLICATION NUMBER: US/09/938,842A
;  CURRENT FILING DATE:  2001-08-24
;  PRIOR APPLICATION NUMBER: US 60/227,866
;  PRIOR FILING DATE: 2000-08-24
;  PRIOR APPLICATION NUMBER: US 60/264,647
;  PRIOR FILING DATE: 2001-01-16
;  PRIOR APPLICATION NUMBER: US 60/300,111
;  PRIOR FILING DATE: 2001-06-22
;  NUMBER OF SEQ ID NOS: 5379
; SEQ ID NO 3346
;   LENGTH: 1881
;   TYPE: DNA
;   ORGANISM: Arabidopsis thaliana
US-09-938-842A-3346

  Query Match             2.9%;  Score 41.4;  DB 9;  Length 1881;
  Best Local Similarity   54.2%;  Pred. No. 1.4;
  Matches   84;  Conservative    0;  Mismatches   71;  Indels    0;  Gaps    0;

Qy    518 ATGGAGTAACTACTGATTATGTTAGATCATTCACAGAATGTGTTAAATGTAGACTTAACT 577
          ||| || | ||   ||| || ||| | | | |||  |  | |||| | ||  ||  ||
Db    537 ATGTAGAATCTTTGGATCATATTATTTTAATGACATTAAATTTTAATTTTAATTTTTTCT 596

Qy    578 TTTACTATAATGGTAATAATGGTAATACTCCTTTCAATCCAGGTAAAAGTTAATGCACAC 637
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Db    597 GTTTTTATTTTGGTGAAATTTGTTTTCTACTTTTTTATATAAGTTATAATAAATGCATAT 656

Qy    638 CTTGTCCGGCAATTAAACCTGCTAATGTTGCTTAA 672
Db    657 TTTGAAAATTAGTTACTTCATTTACAATTTTGTAA 691
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RESULT 15
US-09-864-761-4976/c
; Sequence 4976, Application US/09864761
; Patent No. US20020048763A1
; GENERAL INFORMATION:
; APPLICANT: Penn, Sharron G.
; APPLICANT: Rank, David R.
; APPLICANT: Hanzel, David K.
; APPLICANT: Chen, Wensheng
; TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
; TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
; FILE REFERENCE: Aeomica-X-1
; CURRENT APPLICATION NUMBER: US/09/864,761
; CURRENT FILING DATE: 2001-05-23
; PRIOR APPLICATION NUMBER: US 60/180,312
; PRIOR FILING DATE: 2000-02-04
; PRIOR APPLICATION NUMBER: US 60/207,456
; PRIOR FILING DATE: 2000-05-26
; PRIOR APPLICATION NUMBER: US 09/632,366
; PRIOR FILING DATE: 2000-08-03
; PRIOR APPLICATION NUMBER: GB 24263.6
; PRIOR FILING DATE: 2000-10-04
; PRIOR APPLICATION NUMBER: US 60/236,359
; PRIOR FILING DATE: 2000-09-27
; PRIOR APPLICATION NUMBER: PCT/US01/00666
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00667
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00664
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00669
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00665
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00668
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00663
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00662
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00661
; PRIOR FILING DATE: 2001-01-30
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; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: US 60/234,687
; PRIOR FILING DATE: 2000-09-21
; PRIOR APPLICATION NUMBER: US 09/608,408
; PRIOR FILING DATE: 2000-06-30
; PRIOR APPLICATION NUMBER: US 09/774,203
; PRIOR FILING DATE: 2001-01-29
; NUMBER OF SEQ ID NOS: 49117
; SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
; SEQ ID NO 4976
; LENGTH: 489
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; OTHER INFORMATION: MAP TO AL031076.1
; OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 5.1
; OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 5.5
; OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 5.8
; OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 6
; OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 5.1
; OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 5.2
; OTHER INFORMATION: EXPRESSED IN HBL100, SIGNAL = 4.1
; OTHER INFORMATION: EXPRESSED IN BT474, SIGNAL = 5.9
; OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 5.9
; OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 5.4
US-09-864-761-4976

Query Match 2.9%; Score 41; DB 10; Length 489;
Best Local Similarity 50.2%; Pred. No. 1;
Matches 101; Conservative 0; Mismatches 100; Indels 0; Gaps 0;

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Qy    399 TGCTAGTACATGCACAGCTTGTCCGGTAAACAGAGTTGGTGGTGCATTGACTGCTGGTAA 458
Db    319 TGGTGATTATGGTAGGGATGGTGATGATAATGGTGATGGTGATGGTGAGCATGGTGATGG 260

Qy    459 TGCCGCTACCATAGTCGCATAATGTAACGTCGCATGTCCTACTGGTACTGCACTTGATGA 518
Db    259 TGGTGTTAGTAATCACGGTGATGGTAATGACGGTGGTGGTAATGACGATGATGATGGTGA 200

Qy    519 TGGAGTAACTACTGATTATGT 539
Db    199 TGATGATGGTAATGATAGTAT 179
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Search completed: February 17, 2003, 01:59:09
Job time : 209.08 secs